

Figure 1a

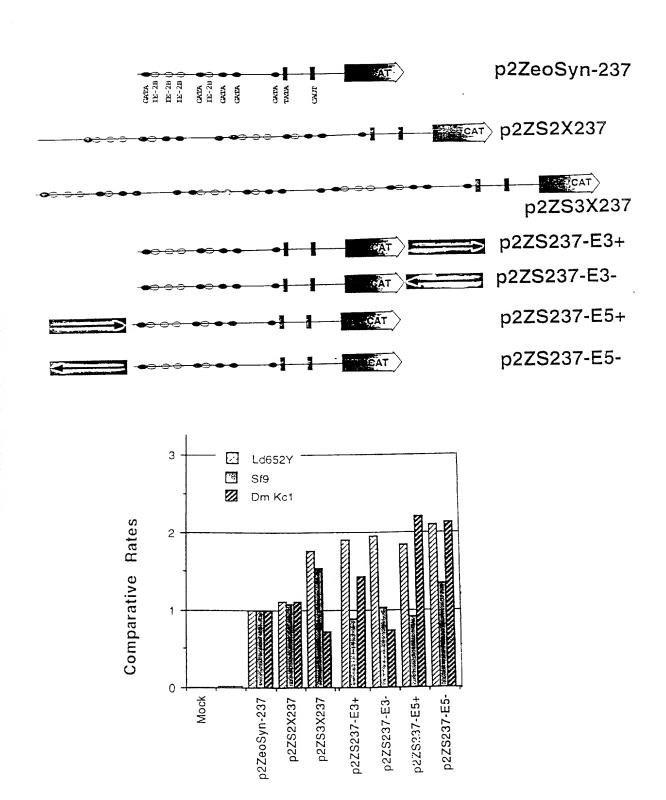


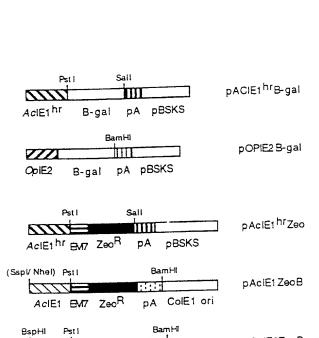
Figure 1b

-375D	-2750	AAATAGICTA CACCGIIGIA TACGCICCAA ATACACTACC ACACATIGAA CCITITITACA GIGCAAAAAA GIACGIGICG GCAGICACGI AGGCCGGC <u>CII</u> 	(a)	(G) (G)	Ageceacece 7	:	30D, -29D -17D  ->5'  IE-2      ->5'       ->		AGC 10C 00
GATAAATTTA							-30D, -29D		(TATA)
E-2		IE-2 IE-N	IE-2	IE-N	IE-2	IE-N		IE-2	IE-N

Figure 2a

-225 -221R -191 -162 -157R -101 -89 -76	
0 000	C
00000	G
0 0 0 <del> </del> 0	U
4444	<
00000	C
00000	G
4 O 4 4 4	4
0000	C
4444	<
0 . U 4 4	1
0 0 1	Ö
0 0 0	G
0 00 0000	$\mathcal{O}$
T TT TTT	L
< << <<<<	4
	H
	H
	$\circ$
0 0	
-232 -212R -197 -175 -147R -118 -93 -80	Consensus
I H G H E D C B A	Cor

## Figure 2b



pOpIE2ZeoB

Figure 3

OpiE2 EM7 ZeoR pA CoiE1 ori

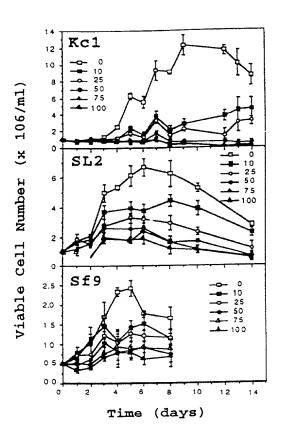


Figure 4a

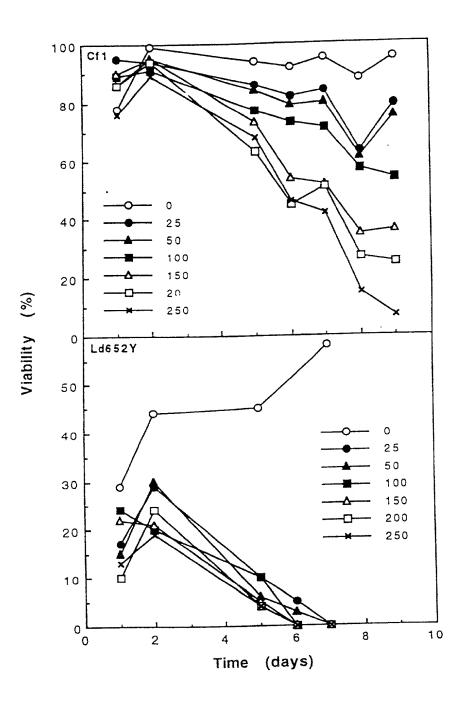


Figure 4b





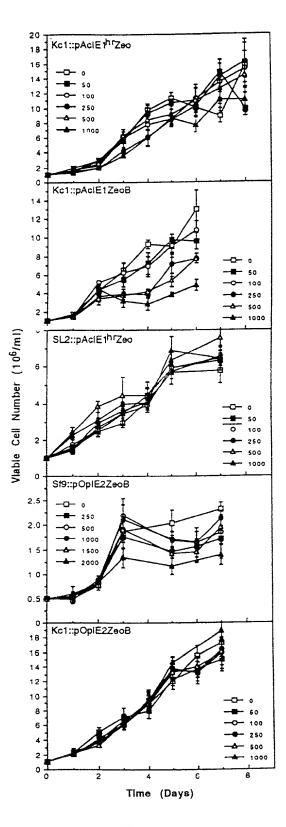


Figure 5

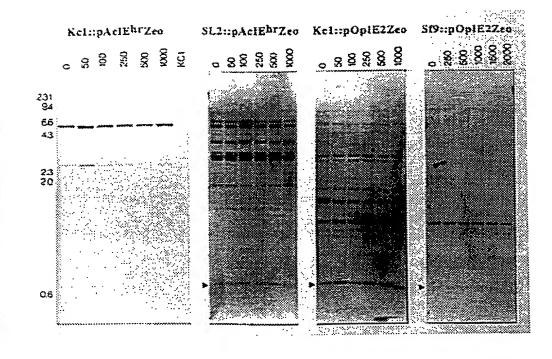


Figure 6

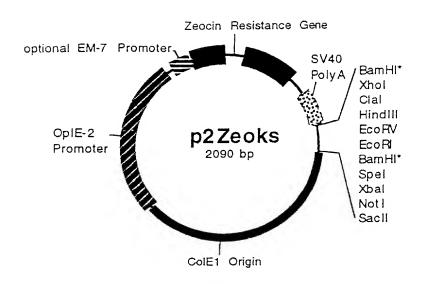


Figure 7

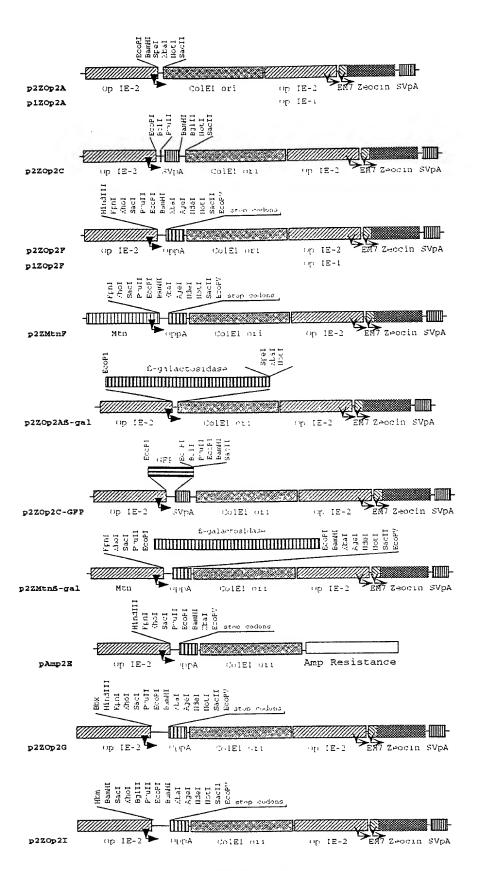


Figure 8a

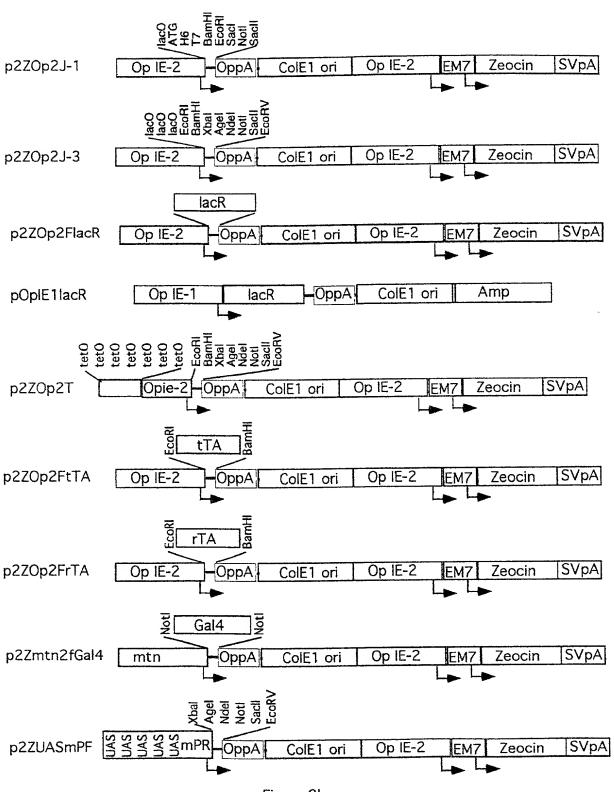
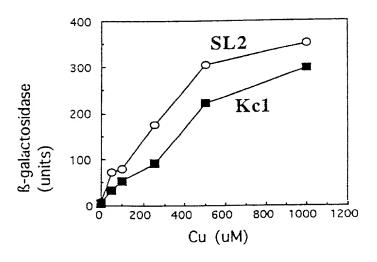


Figure 8b



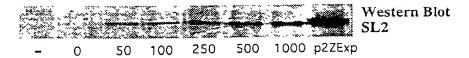


Figure 9

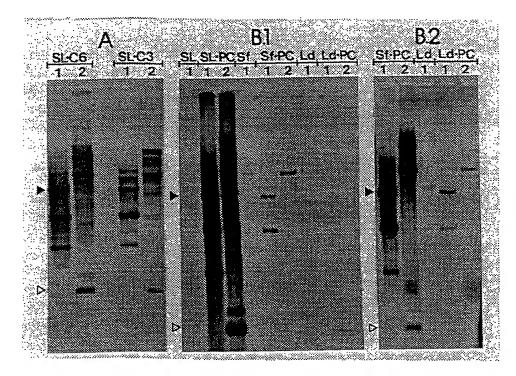


Figure 10

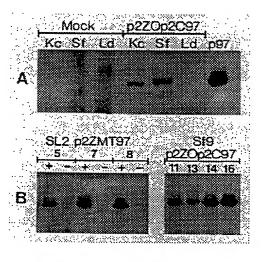


Figure 11





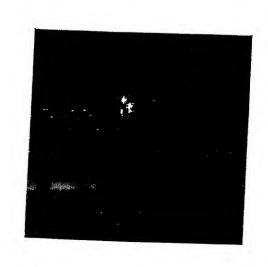


Figure 12 a

CTC TGA TOT CAG CAG TOT TIND SECTENCE OF ON THE CONTROL OF ON THE CONTROL g ., OTG GCG GCC CTG GAA GCC ATG V A A L E G M GAC TAC

*6*8d

ATG ' CTG CAA CCG I 884 8 4 913 > Y Y 9 0 120.7

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GAC TAC OTTS GAC GAC CTS GAA GGS ATTS TEST CAG CAG TAC TAG GAC GAC GAC GAC GAC GAC TAG TAG  $_{
m I}$ 60/90.3

OTE GCS GCS CTB GAA GCS ATTS TCS CCS CCS GCS A A A P S D  $^{\star}$  V A A L E G M S S Q Q C S G A A A P S D  $^{\star}$ Y A g \_ 120.6

TAC OTG GGG GGG ATC TICA Y V A A I \* 9 <u>0</u> 150.5

THE GIE GAT CIE ACT DAA TET TAG Y V D L T K S \* 120.3

chk typ 1 chk typ 2

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Figure 12b

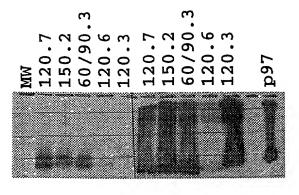


Figure 12c

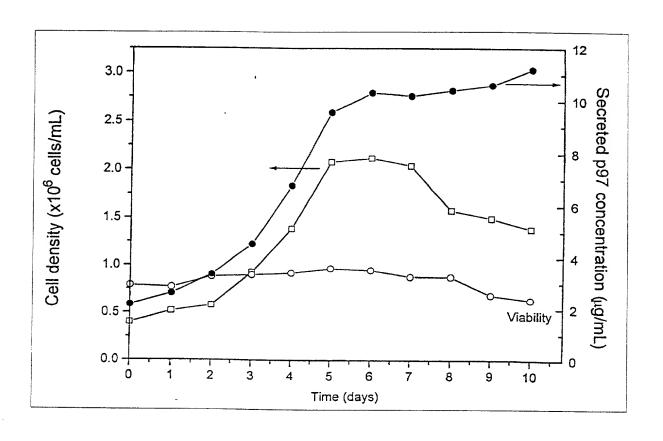


Figure 12d

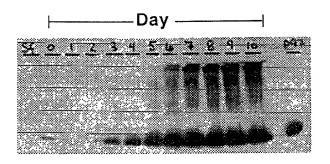


Figure 12e

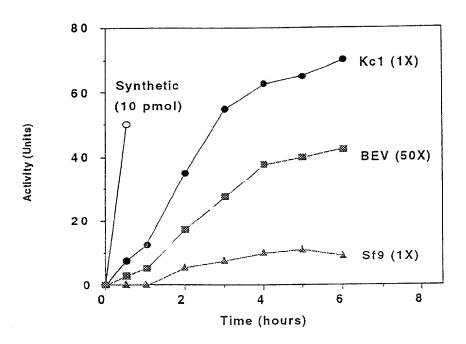


Figure 13a

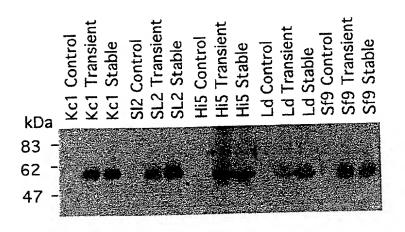


Figure 13b

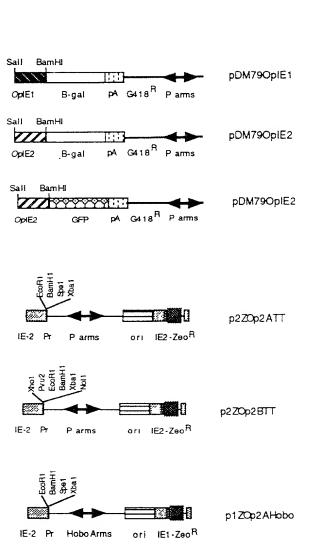


Figure 14

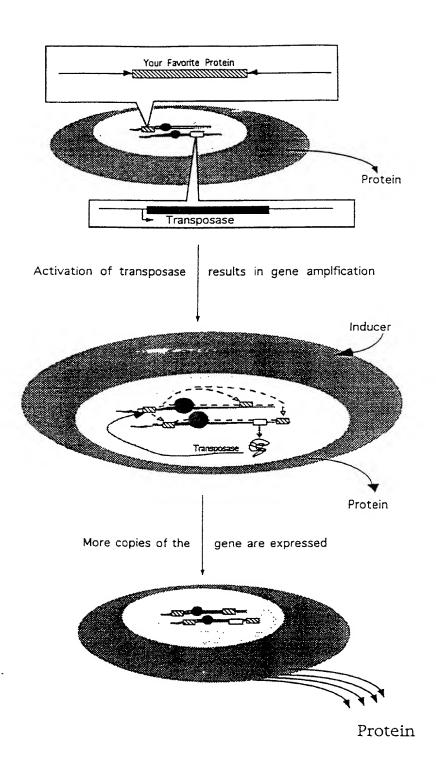


Figure 15

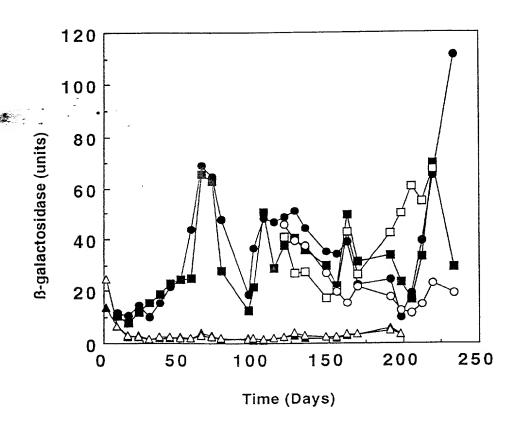


Figure 16a



5' P-element rescue
pDM79  CGACGGGACCACCTTATGTTATTTCATCATGGGCCAGACCCACGTAGTCCAGCGGC
79-2 Xho.6 <b>CGACGGGACCACCTTATGTTATTTCATCATG</b> TCTCGAACCAACGAGAGCAGTATGC
79-2 Xho.4  CGACGGCACCACCTTATGTTATTTCATCATGGTACAGACATCTACTTCCCCCCGCT
79-1 Sph.1  CGACGGGACCACCTTATGTTATTTCATCATGATCTTGCGCTTTAAAATGTGGAGTC
3' P-element rescue
pDM79  CGACGGGACCACCTTATGTTATTTCATCATGGTCTGGCCATTCTCATCGTGAGCTT
79-2 kan.3 <b>CGACGGGACCACCTTATGTTATTTCATCATG</b> AGCCAAACAGAAAGCAGAAAAGCTC
79-2 kan.2 <b>CGACGGGACCACCTTATGTTATTTCATCATG</b> GCCTGACCTAAGCAGATTTGACTGC
79-2 kan.1 CGACGGGACCACCTT CAACGCTACCTAATCTTAAGAACCA
consensus

Figure 16b

GNCYRRAC

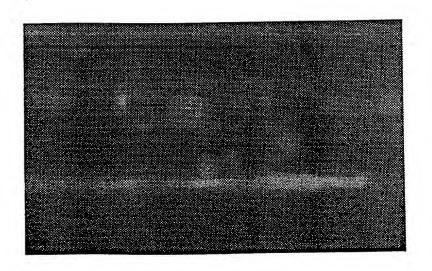


Figure 17